

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/500,601A
P4/10
3/28/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

RAW SEQUENCE LISTING

DATE: 03/28/2005

PATENT APPLICATION: US/10/500,601A

TIME: 14:21:04

Input Set : A:\29808PW1-GE.APP.txt

Output Set: N:\CRF4\03282005\J500601A.raw

2 <110> APPLICANT: Repp Prof., Reinald
 4 <120> TITLE OF INVENTION: Methods for detecting nucleic acids using solid phase
 5 bound primers and cleaving the products of a cyclic
 6 amplification reaction (Trap Release Primer
 7 Amplification (TRamp))
 9 <130> FILE REFERENCE: 29808PWO_RI
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/500,601A
 C--> 12 <141> CURRENT FILING DATE: 2004-06-30
 14 <150> PRIOR APPLICATION NUMBER: DE10164219.9
 15 <151> PRIOR FILING DATE: 2001-12-31
 17 <160> NUMBER OF SEQ ID NOS: 1
 19 <170> SOFTWARE: PatentIn Ver. 2.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 32
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Artificial Sequence
 26 <220> FEATURE:
 27 <221> NAME/KEY: misc_binding
 28 <222> LOCATION: (1)
 29 <223> OTHER INFORMATION: Any (Sequence) ? "n" can only represent a single
 31 <220> FEATURE: nucleotide,
 32 <221> NAME/KEY: misc_binding not a sequence
 33 <222> LOCATION: (32) of nucleotides
 34 <223> OTHER INFORMATION: Any (Sequence) ?
 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: Description of the artificial sequence: PCR-primer
 39 <400> SEQUENCE: 1
 W--> 40 ntgtgtctc agtgttttca ctgagactt tn

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/28/2005
PATENT APPLICATION: US/10/500,601A TIME: 14:21:05

Input Set : A:\29808FW1-GE.APP.txt
Output Set: N:\CRF4\03282005\J500601A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1,32

VERIFICATION SUMMARY

DATE: 03/28/2005

PATENT APPLICATION: US/10/500,601A

TIME: 14:21:05

Input Set : A:\29808PW1-GE.APP.txt

Output Set: N:\CRF4\03282005\J500601A.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:40 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0